



Form 1449 (Modified)	Atty Docket No. MXGNP001X3/124. 610US Applicant: Selifonov et al. Filing Date July 18, 2000	Application No.: 09/618,579 Group 1631
Information Disclosure Statement By Applicant		
(Use Several Sheets if Necessary)		

U.S. Patent Documents

Examiner Initial	No.	Patent No.	Date	Patentee	Class	Sub- class	Filing Date
A	A1	6,537,776 B1	03/25/03	Short	435	69.1	06/14/99
A	A2	6,605,449 B1	08/12/03	Short	435	69.1	06/14/00
	A3						
	A4						
	A5						

Foreign Patent or Published Foreign Patent Application

Examiner Initial	No.	Document No.	Publication Date	Country or Patent Office	Class	Sub- class	Translation	
							Yes	No
	B1							
	B2							
	B3							
	B4							
	B5							

Other Documents

Examiner Initial	No.	Author, Title, Date, Place (e.g. Journal) of Publication
	C1	
	C2	
	C3	
Examiner	Date Considered	

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U.S. Patent Documents

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<i>RA</i>	A1	5,223,409	6/93	Ladner et al.			
	A2	5,198,346	3/93	Ladner et al.			
	A3	6,107,073	8/00	Chen			
	A4	5,741,691	4/21/98	Arnold et al.			
	A5	6,518,065	2/11/03	Stemmer			
	A6	6,365,408	4/2/02	Stemmer			
<i>RA</i>	A7	6,323,030	11/27/01	Stemmer			

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<i>RA</i>	B1	WO03/055978	10/7/03	WIPO			X	

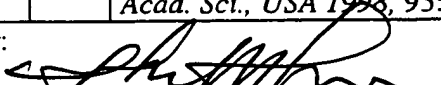
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/	C9	Arnold, "Combinatorial and Computational Challenges for Biocatalyst design", <i>Nature</i> , 2001, 409(6817):253-257
/	C10	Boder et al., "Directed Evolution of Antibody Fragments with Monovalent Femtomolar Antigen binding Affinity", <i>Proc. Natl. Acad. Sci. USA</i> , 2000 97 (20):10701-10705, 2000.
/	C11	Bohm, "New approaches in molecular structure prediction", <i>Biophys Chem.</i> , 1996, 59: 1-32
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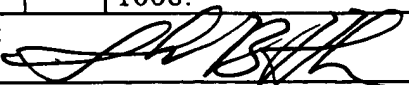
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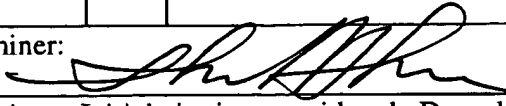
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U.S. Patent Documents

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	A1						
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							Yes	No
	B1							
	B2							
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Other Documents

Examiner Initial	No.	Author, Title, Date, Place (e.g. Journal) of Publication
	C1	CS 262, Computational Genomics, Handout #1: Course Information, printed from: http://www.stanford.edu/class/cs , Spring 2003, 11 pages
A	C2	Corpet et al., Browsing Protein Families Via the Rich Family Description Format," Bioinformatics, Vol. 15, No. 12, 1999, Pages 1020-1027
A	C3	Mironov et al., "Computer Analysis of Transcription Regulatory Patterns in Completely Sequenced Bacterial Genomes," Nucleic Acids Research, Vol. 27, No. 14, 1999, Pages 2981-2989
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